## SEQUENCE LISTING

<110> HINUMA, Shuji KOBAYASHI, Makoto HABATA, Yugo HARADA, Masataka OKUBO, Shoichi YOSHIDA, Hiromi NISHI, Kazunori

<120> A Novel Ligand For FPRL1 And Its Use

<130> 3118 USOP

<140> US/10/534,082

<141> 2005-05-05

<150> PCT/JP2003/014138

<151> 2003-11-06

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<151> 2002-11-07

<150> JP 2002-367119

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<150> JP 2003-59073

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<213> Homo sapiens

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Arg	Val	Leu	Thr	Ala 245	Val	Val	Ala	Ser	Phe 250	Phe	Ile	Cys	Trp	Phe 255	Pro	
Phe	Gln	Leu	Val 260	Ala	Leu	Leu	Gly	Thr 265	Val	Trp	Leu	Lys	Glu 270	Met	Leu	•
Phe	Tyr	Gly 275	Lys	Tyr	Lys	Ile	Ile 280	Asp	Ile	Leu	Val	Asn 285	Pro	Thr	Ser	
Ser	Leu 290	Ala	Phe	Phe	Asn	Ser 295	Cys	Leu	Asn	Pro	Met 300	Leu	Tyr	Val	Phe	
Val 305	Gly	Gln	Asp	Phe	Arg 310	Glu	Arg	Leu	Ile	His 315	Ser	Leu	Pro	Thr	Ser 320	
Leu	Glu	Arg	Ala	Leu 325	Ser	Glu	Asp	Ser	Ala 330	Pro	Thr	Asn	Asp	Thr 335	Ala	
Ala	Asn	Ser	Ala 340	Ser	Pro	Pro	Ala	Glu 345	Thr	Glu	Leu	Gln	Ala 350	Met	• ,	
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cta	aac	cta	acc	cta	act	gac	+++	tct	ttc	асσ	acc	aca	tta	cca	ttc	240

Leu 65	Asn	Leu	Ala	Leu	Ala 70	Asp	Phe	Ser	Phe	Thr 75	Ala	Thr	Leu	Pro	Phe 80		•
				_	gcc Ala	-		-									288
		-			cac His				-						-	• .	336
					ttc Phe												384
					cag Gln												432
					tgg Trp 150												480
					aca Thr												528
					tcc Ser												576
				-	ctg Leu												624
		-	_		atg Met						-						672
_	_	_			aaa Lys 230	_		_				_	_				720
					gtg Val												768
					ctt Leu												816
					aaa Lys												864
					aac Asn												912

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	g gag ı Glu									Pro						1008
	aat Asn															1053
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Туі	Asp	Ser	Thr 20	Ile	Ser	Arg	Val	Leu 25	Trp	Ile	Leu	Thr	Met 30	Val	Val	
Let	ı Ser	Ile 35	Thr	Phe	Val	Leu	Gly 40	Val	Leu	Gly	Asn	Gly 45	Leu	Val	Ile	
	Val 50		_		_	55					60			· -		
Let 65	Asn	Leu	Ala	Leu	Ala 70	Asp	Phe	Ser	Phe	Thr 75		Thr	Leu	Pro	Phe 80	
Phe	e Val	Ile	Ser	Ile 85	Ala	Met	Lys	Glu	Lys 90	Trp	Pro	Phe	Gly	Trp 95	Phe	
	ı Cys	•	100					105					110	* 7		
	. Phe	115					120					125		*		••
	130					135	,			,	140					
Va]	. Val	Val	Gly	Pro	Trp 150	Ile	Leu	Ala		Ile 155	Leu	Thr	Leu	Pro		
	: Ile	Phe	Met	Thr		Val	Arg	Ile			Glv	Asn	Val	Tvr	160 Cvs	•.
	Phe			165					170	*				175		
	. Ala		180					185					190	•		
	Phe	195					200			•		205				
	210 Val				•	215					220	-	_			
225		,			23.0					235					240	
	Gln			245					250			_	_	255		
			260					265					270			٠ .
	Ser	275					280					285				
sei	Leu	AId	TÄT	rne	ASII	ser.	cys	ьeu	ASII	Pro	Mer	ьeu	LAI.	AIG	rne	

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290
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                                       315
Leu Glu Arg Ala Leu Ser Glu Asp Ser Gly Gln Thr Ser Asp Thr Gly
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                                                                   120
gtgctgggta atggactagt gatctgggta gctggattcc ggatggtaca cactgtcacc
                                                                   180
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tttgtcatct caattgctat gaaagaaaaa tggccttttg gatggttcct gtgtaaatta
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gttcacattg tagtagacat aaacctcttt ggaagtgtct tcctgattgc tttaattgcc
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                                                                   420
ctggctagga aggtggttgt tgggccctgg attttagctc tgattctcac tttgcccatt
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gcatcctggg gtaacactgc tgaagaacta ttgaacatag ctaacacttt tgtaacagtt
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                                                                   840
                                                                   900
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ctggagagag ccctgagtga ggactctggc caaaccagtg atacaggcat cagttctgct
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Trp Val Ala Gly Phe Arg Met Pro His Thr Val Thr Thr Ile Trp Tyr
Leu Asn Leu Ala Leu Ala Asp Phe Ser Phe Thr Ala Thr Leu Pro Phe
                    70
                                        75
Leu Leu Val Glu Met Ala Met Lys Glu Lys Trp Pro Phe Gly Trp Phe
                                    90
Leu Cys Lys Leu Val His Ile Val Val Asp Val Asn Leu Phe Gly Ser
                               105
Val Phe Leu Ile Ala Leu Ile Ala Leu Asp Arg Cys Ile Cys Val Leu
       115
                           120
                                               125
His Pro Val Trp Ala Gln Asn His Arg Thr Val Ser Leu Ala Arg Lys
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140

135

130

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Phe Ile Phe Leu Thr Thr Val Arg Ile Pro Gly Gly Asp Val Tyr Cys
                                     170
Thr Phe Asn Phe Gly Ser Trp Ala Gln Thr Asp Glu Glu Lys Leu Asn
                                 185
Thr Ala Ile Thr Phe Val Thr Thr Arg Gly Ile Ile Arg Phe Leu Ile
                            200
                                                 205
Gly Phe Ser Met Pro Met Ser Ile Val Ala Val Cys Tyr Gly Leu Ile
                        215
Ala Val Lys Ile Asn Arg Arg Asn Leu Val Asn Ser Ser Arg Pro Leu
                    230
                                        235
Arg Val Leu Thr Ala Val Val Ala Ser Phe Phe Ile Cys Trp Phe Pro
                245
                                    250
Phe Gln Leu Val Ala Leu Leu Gly Thr Val Trp Phe Lys Glu Thr Leu
            260
                                265
Leu Ser Gly Ser Tyr Lys Ile Leu Asp Met Phe Val Asn Pro Thr Ser
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Ser Leu Ala Tyr Phe Asn Ser Cys Leu Asn Pro Met Leu Tyr Val Phe
                        295
Met Gly Gln Asp Phe Arg Glu Arg Phe Ile His Ser Leu Pro Tyr Ser
                    310
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                                                                     180
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                                                                     240
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                                                                     780
gcccttttgg gcacagtctg gtttaaagag acattgctta gtggtagtta taaaattctt
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                                                                     900
ctctatgttt tcatgggcca ggactttcgt gagagattta ttcattccct gccttatagt
                                                                     960
cttgagagag ccctgagtga ggattctggt caaaccagtg attcaagcac cagttctact
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<223> Primer

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